## CLUSTAL W(1.60) multiple sequence alignment

Haar D. Com	
UserP: Hithat 1991 4 UserP: Market 1991 4 UserP: 1994 999	HACSNPAPQASRAAAERVPPLLAARPCTPARSRSGPPPPQRRLLLLFLSPSLSVSRSLSP
	SRFSSSSTVACPGE GRARP VCW LQPSRGASSSMSSDMAADESEAPVLSEDEVWEFCLDKTEDGGGSPGSDVTDTCEPPCGCW MFRLRJWASSTTGSRYGS AFCGSP *
UserP: Monso (FDD 4   1   UserP: Monso (FDD 4   1   UserP: MGGBP   1	KRS EMATCAVEVFGLLEDEENSR1 ELNPNSLEEEHVLFTADPYLELHN DDTKV TLAWCYCVPVCYGES F1
UserP: Morse NEDD 4 UserP: Morse NEDD 4 UserP: GG3P	VP
UserP: Mouse NEDD 4 I	LYPLPTENPRLERPYTFFDFVLHPRSHKSFVKGYLRLKMTYLPKTSGSEDDNAEQAEELE LYPLPTENPRMERPYTFKDFVLHPRSHKSFVKGYLRLKMTYLPKN GSEDENADQAEELE LSHLPTEDPTMEFPYTFKDFLLRPRSHKSFVKGFLRLKMAYMPKNGGQDEENSDQRDDME
UserP:Mouse_NEDD 4 I	PGWVVLDQPDAACHLQQQQEP PGWVVLDQPDAATHLPHPPEP DDDLT HGWEVVDSNDSASQHQEELPPPP SLMDVSS
UserP: Mouse NEDD 4 I UserP: MGGBP I	DAFNGNIÇLQAQRAFTTRRQISEETES -VDNQESSENWEIIREDEATMYSSQAFPS DEDNDDMQLQAQRAFTTRRQISEDVDGPDNRESPENWEIVREDENTEYSGQAVQS ESDNNIRQINQEAAHRRFRSRRHISEDLFPEPSEGGEVPEPWETISEEVNIAGDSLGVVL
UserP:Mouse NEDD 4 I	PPPSSNLDVPTHLAEELNARLTIFGNSAVSQPASSSN HSSF RGSLQAYTFEEQP PPSGHIDVQTHLAEEFNTRLAVCGNPATSQPVTSSN HSSF RGSLQAYTFEEQP PPPPASFGSRTSPQELSEELSRRLQITPDSNGEQFSSLIQPEPSSPLRSCSVTDAVAEQG *** *** *** *** *** *** *** *** *** **
UserP:Mouse NEDD 4 7	TUP VLLPTS QATVETSQLTSQ TUP VLLPTS Q HLPPPSVAYVHTTPG IMQLAEDGASGSATN
UserP: Human NEDD 4 UserP: Mouse NEDD 4 UserP: DGG9P 5	SS AGPQSQA STSDS GQQVTQPS D  SIDHLIEPQIFRFRSLSSPTVTLXAPLEGAKDSPVRFAVKLTLSNFQSPQPSPY::SPKPQ
UserP: Human NEDD 4 UserP: Monso NEDD 4 UserP: GGGRP F	EIEQ FIPAHLPGFTSLDTSMDL  DFFSFIPAHLFGFTD SMDL  HKNTQSF FLEFPVHMPSETSLN PMDLGP
UserP: Montage NEDD 4 UserP: Montage NEDD 4 UserP: MGG39	ENVAITGPAVPYSPDYKRKYEFFREKLEK QNVAITGPAVPYSPDYKRKYEFFHEKLEK PLQNPAITGPAVPYSPEFFQKYDYFFKELEK
UserP: Mariago NFDD 4 (	QMI PNY FEMFLERATVLEDSYRFIMGVFFADFLFAFLWIEFDGEFGLDYGGVAFEWFFL QMI PNY FEMFLERANILEDSYRFIMGVFFADILFAFLWIEFDGEFGLDYGGVAFEWFFL FALIPNFFEMFLHRINIFEESYRFIMSVFFPDVLFARLWIEFESEFGLDYGGVAFEWF
	ISFEMFNPYYGLFEYSATDNYTL@INPNSGLCNEDHLSYFFFIGFVAGMAVYHGFLLDGF ISFEMFNPYYGLFEYSATDNYTL@INPNSGLCNEDHLSYFFFIGFVAGMAVYHGFLLDGF

UserP: <u>13433 ]</u>	
	FIRPFYKMMLHKPITLHDMESVDSEYYNSLRWILENDPTELDLRFIIDEELFGQTHQHEL FIRPFYKMMLQKLITLHDMESVDSEYYSSLRWILENDPTELDLRFIIDEELFGQTHQHEL
	KNGGSEIVVTNKNKKEYIYLVIQWRFVNRIQKQMAAFKEGFFELIPQDLIKIFDENELEL KTGGSEIVVTNKNKKEYIYLVIQWRFVNRIQKQMAAFKEGFFELIPQDLIKIFDENELEL
	LMCGLGDVDVNDWREHTK LMCGLGDVDVNDWREHTK
UserP:Human NEDD 4 UserP:Mouse NEDD 4 UserP:76GBP 1	**************************************
UserP:Himail NEDD 4 UserP:Mous+ NEDD 4 UserP: <u>TGGBP</u> 1	DX * _ * * *

Exhibit 3: Clustal W alignment of ZGGBP-1 to the human and mouse forms of NEDD-4. The functional domains identified using Hmm for ZGGBP-1 are shown in Red. The equivalent functional domains annotated in SwissProt for the human and mouse NEDD-4 are shown in green.